Wyoming-Specific Activity: MMWR Week 11 (Week ending March 21, 2009)

Week	Total			
40	8			
41	4			
42	0			
43	2			
44	0			
45	1			
46	3			
47	1			
48	0			
49	1			
50	0			
51	1			
52	2			
53	1			
1	2			
2	1			
3	7			
4	20			
5	39			
6	65			
7	74			
8	107			
9	129			
10	109			
11	135			
12				
13				
14				
15				
16				
17				
18				
19				
20				
Unknown				
Total	713			

County	Totals				
Albany	37 [*]				
Big Horn	20				
Campbell	55				
Carbon					
Converse	8				
Crook	6				
Fremont	41				
Goshen	7				
Hot Springs	6				
Johnson					
Laramie	299				
Lincoln	8^*				
Natrona	94				
Niobrara	20*				
Park	/ ()				
Platte	9*				
Sheridan	6*				
Sublette	29				
Sweetwater	31				
Teton	14				
Uinta	5				
Washakie	8				
Weston	8				
Unknown					
Total	713				

Age	Number
0-4	149
5-10	148
11-19	156
20-39	158
40-59	74
60+	28
Unknown	
Total	713

Gender	Number			
Male	366			
Female	347			
Unknown				
Total	713			

Type	Number			
A	384			
В	171			
Unknown	158			
Total	713			

Test	Number		
Rapid	699		
Culture	11		
PCR	1		
DFA	1		
IFA	1		
Total	713		

^{*} Counties with positive laboratory cultures

Wyoming Public Health Laboratory Testing: MMWR Week 11 (Week ending March 21, 2009)

Week	# Submitted	A (H1)	A (H3)	В	Negative	Unknown	Not Tested
40	1	-	-	-	1		
41	0	-	-	-	-		
42	0	-	-	-	-		
43	0	-	-	-	-		
44	1	-	-	-	1		
45	0	Ī	-	-	-		
46	0	-	-	-	-		
47	2	-	-	-	2		
48	0	-	-	-	-		
49	1	-	-	-	1		
50	1	-	-	-	1		
51	0	-	-	-	-		
52	0	Ī	-	-	-		
53	0	1	-	-	-		
1	0	-	-	-	-		
2	0	-	-	-	-		
3	2	1	1	-	-		
4	4	-	-	1	3		
5	4	-	2	-	2		
6	1	-	-	-	1		
7	1	-	1	-	-		
8	3	Ī	1	1	1		
9	1	1	-	-	1		
10	6	1	1	-	4		
11	4	1	-	1	3		
12							
13							
14							
15							
16							
17							
18							
19							
20							
Total	32	2	6	3	21	0	0

Antigenic Characterization: MMWR Week 11 (Week ending March 21, 2009)

The Centers for Disease Control and Prevention (CDC) has antigenically characterized 807 influenza viruses [510 influenza A (H1), 86 influenza A (H3) and 211 influenza B viruses] collected by U.S. laboratories since October 1, 2008.

All 510 influenza A (H1) viruses are related to the influenza A (H1N1) component of the 2008-09 influenza vaccine (A/Brisbane/59/2007). All 86 influenza A (H3N2) viruses are related to the A (H3N2) vaccine component (A/Brisbane/10/2007).

Influenza B viruses currently circulating can be divided into two distinct lineages represented by the B/Yamagata/16/88 and B/Victoria/02/87 viruses. Forty-four (20.9%) influenza B viruses tested belong to the B/Yamagata lineage and are related to the vaccine strain (B/Florida/04/2006). The remaining 167 (79.1%) viruses belong to the B/Victoria lineage and are not related to the vaccine strain.

Data on antigenic characterization should be interpreted with caution given that antigenic characterization data is based on hemagglutination inhibition (HI) testing using a panel of reference ferret antisera and results may not correlate with clinical protection against circulating viruses provided by influenza vaccination.

Annual influenza vaccination is expected to provide the best protection against those virus strains that are related to the vaccine strains, but limited to no protection may be expected when the vaccine and circulating virus strains are so different as to be from different lineages, as is seen with the two lineages of influenza B viruses.